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**EP 0 318 216 B1**

sequence of the polypeptide encoded in the extended ORF in the derived sequence.

Fig. 27 shows the sequence of the HCV cDNA in clone 12f, the segment which overlaps clone 14i, and the amino acids encoded therein.

Fig. 28 shows the sequence of the HCV cDNA in clone 35f, the segment which overlaps clone 39c, and the amino acids encoded therein.

Fig. 29 shows the sequence of the HCV cDNA in clone 19g, the segment which overlaps clone 35f, and the amino acids encoded therein.

Fig. 30 shows the sequence of clone 26g, the segment which overlaps clone 19g, and the amino acids encoded therein.

Fig. 31 shows the sequence of clone 15e, the segment which overlaps clone 26g, and the amino acids encoded therein.

Fig. 32 shows the sequence in a composite cDNA, which was derived by aligning clones 12f through 15e in the 5' to 3' direction; it also shows the amino acids encoded in the continuous ORF.

Fig. 33 shows a photograph of Western blots of a fusion protein, SOD-NANB<sub>5-1-1</sub>, with chimpanzee serum from chimpanzees infected with BB-NANB, HAV, and HBV.

Fig. 34 shows a photograph of Western blots of a fusion protein, SOD-NANB<sub>5-1-1</sub>, with serum from humans infected with NANBV, HAV, HBV, and from control humans.

Fig. 35 is a map showing the significant features of the vector pAB24.

Fig. 36 shows the putative amino acid sequence of the carboxy-terminus of the fusion polypeptide C100-3 and the nucleotide sequence encoding it.

Fig. 37A is a photograph of a coomassie blue stained polyacrylamide gel which identifies C100-3 expressed in yeast.

Fig. 37B shows a Western blot of C100-3 with serum from a NANBV infected human.

Fig. 38 shows an autoradiograph of a Northern blot of RNA isolated from the liver of a BB-NANBV infected chimpanzee, probed with BB-NANBV cDNA of clone 81.

Fig. 39 shows an autoradiograph of NANBV nucleic acid treated with RNase A or DNase I, and probed with BB-NANBV cDNA of clone 81.

Fig. 40 shows an autoradiograph of nucleic acids extracted from NANBV particles captured from infected plasma with anti-NANB<sub>5-1-1</sub>, and probed with <sup>32</sup>P-labeled NANBV cDNA from clone 81.

Fig. 41a and b shows autoradiographs of filters containing isolated NANBV nucleic acids, probed with <sup>32</sup>P-labeled plus and minus strand DNA probes derived from NANBV cDNA in clone 81.

Fig. 41-1 shows the homologies between a polypeptide encoded in HCV cDNA and an NS protein from Dengue flavivirus.

Fig. 43 shows a histogram of the distribution of HCV infection in random samples, as determined by an ELISA screening.

Fig. 44 shows a histogram of the distribution of HCV infection in random samples using two configurations of immunoglobulin-enzyme conjugate in an ELISA assay.

Fig. 45 shows the sequences in a primer mix, derived from a conserved sequence in NS1 of flaviviruses.

Fig. 46 shows the HCV cDNA sequence in clone k9-1, the segment which overlaps the cDNA in Fig. 27, and the amino acids encoded therein.

Fig. 47 shows the sequence in a composite CDNA which was derived by aligning clones k9-1 through 15e in the 5' to 3' direction; it also shows the amino acids encoded in the continuous ORF.

#### I. Definitions

The term "hepatitis C virus" has been reserved by workers in the field for an heretofore unknown etiologic agent of NANBH. Accordingly, as used herein, "hepatitis C virus" (HCV) refers to an agent causative of NANBH, which agent is a virus characterised by: (i) a positive stranded RNA genome; (ii) said genome comprising an open reading frame (ORF) encoding a polyprotein; and (iii) the portion of said polyprotein corresponding to Figure 14 having at least 40% homology to the amino acid sequence in Figure 14. This agent was formerly referred to as NANBV and/or BB-NANBV. The terms HCV, NANBV, and BB-NANBV are used interchangeably herein, but all refer to the virus as defined above. As an extension of this terminology, the disease caused by HCV, formerly called NANB hepatitis (NANBH), is called hepatitis C. The terms NANBH and hepatitis C may be used interchangeably herein.

The term "HCV", as used herein, denotes a viral species which causes NANBH, and attenuated strains or defective interfering particles derived therefrom. As shown infra., the HCV genome is comprised of RNA. It is known that RNA containing viruses have relatively high rates of spontaneous mutation, i.e., reportedly

FIG. 32-1 COMBINED ORF OF DNAs 12f through 15e

IlePheLysIleArgMetTyrValGlyGlyValGluHisArgLeuGluAlaAlaCysAsn  
 1 CCATATTTAAATCAGGATGTACGTGGGAGGGGTCGAACACAGGCTGGAAGCTGCCTGCA  
 GGTATAAATTTTAGTCTACATGCACCTCCCGAGCTTGTGTCGACCTTCGACGGACGT  
 TrpThrArgGlyGluArgCysAspLeuGluAspArgAspArgSerGluLeuSerProLeu  
 61 ACTGGACGCGGGCGAACGTTGCGATCTGGAAGACAGGGACAGGTCCGAGCTCAGCCCGT  
 TGACCTGCGCCCCGCTTGCAACGCTAGACCTTCTGTCCCTGTCCAGGCTCGAGTCGGGCA  
 LeuLeuThrThrThrGlnTrpGlnValLeuProCysSerPheThrThrLeuProAlaLeu  
 121 TACTGTGACCACTACACAGTGGCAGGTCTCCCGTGTTCCTTCACAACCTACCAGCCT  
 ATGACGACTGGTGATGTGTACCGTCCAGGAGGGCACAAGGAAGTGTGGGATGGTCGGA  
 SerThrGlyLeuIleHisLeuHisGlnAsnIleValAspValGlnTyrLeuTyrGlyVal  
 181 TGTCCACCGGCCTCATCCACTCCACCAGAACATTGTGGACGTGCAGTACTTGTACGGGG  
 ACAGGTGGCCGGAGTAGGTGGAGGTGGTCTTGTAAACCTGCACGTATGAACATGCCCC  
 GlySerSerIleAlaSerTrpAlaIleLysTrpGluTyrValValLeuLeuPheLeuLeu  
 241 TGGGGTCAAGCATCGCGTCTGGGCCATTAGTGGGAGTACGTGCTTCCTGTTCCTTC  
 ACCCCAGTTTCGTAGCGCAGGACCGGTAATTACCTCATGCAGCAAGAGGACAAGGAAG  
 LeuAlaAspAlaArgValCysSerCysLeuTrpMetMetLeuLeuIleSerGlnAlaGlu  
 301 TGGTTCGAGACGCGCGCTCTGCTCCTGCTTGTGGATGATGCTACTCATATCCCAAGCGG  
 ACGAACGTCTGCGCGCGCAGACGAGGACGAACACCTACTACGATGAGTATAGGGTTCGCC  
 AlaAlaLeuGluAsnLeuValIleLeuAsnAlaAlaSerLeuAlaGlyThrHisGlyLeu  
 361 AGGCGGCTTTGGAGAACCTCGTAATACTTAATGCAGCATCCCTGGCCGGGACGCAAGGTC  
 TCCGCCGAAACCTCTTGGAGCATTATGAATTACGTGCTAGGGACCGGCCCTGCGTGCCAG  
 ValSerPheLeuValPhePheCysPheAlaTrpTyrLeuLysGlyLysTrpValProGly  
 421 TTGTATCTTCTCTGTTCTTCTGCTTTCATGTTATTTGAAGGGTAAGTGGGTGCCCCG  
 AACATAGGAAGGAGCACAAGAAGACGAAACGTACCATAAACTTCCATTACCCACGGGC  
 AlaValTyrThrPheTyrGlyMetTrpProLeuLeuLeuLeuLeuAlaLeuProGln  
 481 GAGCGGTCTACACCTTCTACGGGATGTGGCCTCTCTCTCTGCTCTGTTGGCGTTGCCCC  
 CTCGCCAGATGTGGAAGATGCCCTACACGGGAGAGGAGGACGAGGACAACCGCAACGGGG  
 ArgAlaTyrAlaLeuAspThrGluValAlaAlaSerCysGlyGlyValValLeuValGly  
 541 AGCGGGCGTACGCGCTGGACACGGAGGTGGCCGCGTCTGTGGCGGTGTTGTTCTCGTCG  
 TCGCCCGCATGCGCGACCTGTGCCTCCACCGGCGCAGCACACGCCACAACAAGAGCAGC  
 LeuMetAlaLeuThrLeuSerProTyrTyrLysArgTyrIleSerTrpCysLeuTrpTrp  
 601 GGTGTATGGCGCTGACTCTGTACCATATTACAAGCGCTATATCAGCTGGTGCTTGTGGT  
 CCAACTACCGCGACTGAGACAGTGGTATAATGTTCCGATATAGTCGACCAACGAACCA  
 LeuGlnTyrPheLeuThrArgValGluAlaGlnLeuHisValTrpIleProProLeuAsn  
 661 GGCTTCAGTATTTTCTGACCAGAGTGAAGCGCAACTGCACGTGTGGATTCCCCCCTCA  
 CCGAAGTCATAAAAGACTGGTCTCACCTTCGCGTTGACGTGCACACCTAAGGGGGGGAGT  
 ValArgGlyGlyArgAspAlaValIleLeuLeuMetCysAlaValHisProThrLeuVal  
 721 ACGTCCGAGGGGGGCGCGACCGCTCATCTTACTCATGTGTGCTGTACACCGACTCTGG  
 TGCAGGCTCCCCCGCGCTGCGGCAGTAGAATGAGTACACACGACATGTGGGCTGAGACC  
 PheAspIleThrLysLeuLeuAlaValPheGlyProLeuTrpIleLeuGlnAlaSer  
 781 TATTTGACATCACCAATGTCTGCTGGCGCTTTCGGACCCCTTTGGATTCTTCAAGCCA  
 ATAACTGTAGTGGTTTAAACGACGACCGGCGAGAAGCCTGGGGAAACCTAAGAAGTTCGGT  
 LeuLeuLysValProTyrPheValArgValGlnGlyLeuLeuArgPheCysAlaLeuAla  
 841 GTTGTCTAAAGTACCCTACTTGTGCGCGTCCAAGGCCTTCTCGGTTCTGCGCGTTAG  
 CAAACGAATTTTCATGGGATGAAACACGCGCAGGTTCCGGAAGAGGCCAAGACGCGCAATC  
 ArgLysMetIleGlyGlyHisTyrValGlnMetValIleIleLysLeuGlyAlaLeuThr  
 901 CGCGGAAGATGATCGGAGGCCATTACGTGCAAATGGTTCATCATTAAAGTTAGGGGCGCTTA  
 CGCCTTCTACTAGCCTCCGGTAATGCACGTTTACCAGTAGTAATTCAATCCCCGCGAAT

- 961 GlyThrTyrValTyrAsnHisLeuThrProLeuArgAspTrpAlaHisAsnGlyLeuArg  
CTGGCACCTATGTTTATAACCATCTCACTCCTCTTCGGGACTGGGCGCACAAACGGCTTGC  
GACCGTGGATACAAATATTGGTAGAGTGAGGAGAAGCCCTGACCGCGTGTGCGGAACG
- 1021 AspLeuAlaValAlaValGluProValValPheSerGlnMetGluThrLysLeuIleThr  
GAGATCTGGCCGTGGCTGTAGAGCCAGTCGTCTTCTCCCAAATGGAGACCAAGCTCATCA  
CTCTAGACCGGCACCGACATCTCGGTACGACAGAGAGGGTTTACCTCTGGTTCGAGTAGT
- 1081 TrpGlyAlaAspThrAlaAlaCysGlyAspIleIleAsnGlyLeuProValSerAlaArg  
CGTGGGGGGCAGATACCGCCCGTGGGTGACATCATCAACGGCTTGCCTGTTTCCGCCC  
GCACCCCCCGTCTATGGCGGCGACGCCACTGTAGTAGTTGCCGAACGGACAAAGGCGGG
- 1141 ArgGlyArgGluIleLeuLeuGlyProAlaAspGlyMetValSerLysGlyTrpArgLeu  
GCAGGGGCGGGAGATACTGCTCGGGCCAGCCGATGGAATGGTCTCCAAGGGGTGGAGGT  
CTCCCCCGGCCCTCTATGACGAGCCCGTGGCTACCTTACCAGAGGTTCCCCACCTCCA
- 1201 LeuAlaProIleThrAlaTyrAlaGlnGlnThrArgGlyLeuLeuGlyCysIleIleThr  
TGCTGGCGCCCATCACGGCGTACGCCAGCAGACAAGGGGCTCTAGGGTGCTAATCA  
ACGACCGCGGGTAGTGCGCATGCGGGTCTGTTCCCGGAGGATCCACGTATTAGT
- 1261 SerLeuThrGlyArgAspLysAsnGlnValGluGlyGluValGlnIleValSerThrAla  
CCAGCCTAACTGGCCGGGACAAAACCAAGTGGAGGGTGAGGTCCAGATTGTGTCAACTG  
GGTCGGATTGACCGGCCCTGTTTGGTTTCACTCCACTCCAGGTCTAACACAGTTGAC
- 1321 AlaGlnThrPheLeuAlaThrCysIleAsnGlyValCysTrpThrValTyrHisGlyAla  
CTGCCCAAACCTTCTGGCAACGTGCATCAATGGGGTGTGCTGGACTGTCTACCACGGGG  
GACGGGTTTGGAAGGACCGTTGCACGTAGTTACCCACACGACCTGACAGATGGTGCCCC
- 1381 GlyThrArgThrIleAlaSerProLysGlyProValIleGlnMetTyrThrAsnValAsp  
CCGGAACGAGGACCATCGCGTACCCAAGGGTCTGTTCATCCAGATGTATACCAATGTAG  
GGCCTTGCTCCTGGTAGCGCAGTGGGTTCCAGGACAGTAGGTCTACATATGGTTACATC
- 1441 GlnAspLeuValGlyTrpProAlaProGlnGlySerArgSerLeuThrProCysThrCys  
ACCAAGACCTTGTGGGCTGGCCCGCTCCGCAAGGTAGCCGCTCATTGACACCCTGCACCT  
TGGTTCTGGAACACCGACCGGGCGAGGCGTTCCATCGGCGAGTAACTGTGGGACGTGAA
- 1501 GlySerSerAspLeuTyrLeuValThrArgHisAlaAspValIleProValArgArgArg  
GCGGCTCCTCGGACCTTTACCTGGTACGAGGCACGCCGATGTCTATCCCGTGGCGCGGC  
CGCGAGGAGCCTGGAAATGGACCACTGCTCCGTGGGCTACAGTAAGGGCACGCGGGCG
- 1561 GlyAspSerArgGlySerLeuLeuSerProArgProIleSerTyrLeuLysGlySerSer  
GGGGTGATAGCAGGGGAGCCTGCTGTGCGCCCGGCCCATTTCTACTTGAAAGGCTCCT  
CCCCACTATCGTCCCCGTGGACGACAGCGGGGCGGGTAAAGGATGAACCTTCGAGGA
- 1621 GlyGlyProLeuLeuCysProAlaGlyHisAlaValGlyIlePheArgAlaAlaValCys  
CGGGGGGTCCGCTGTTGTGCCCCGCGGGGCACGCCGTGGGCATATTTAGGGCCGCGGT  
GCCCCCAGGCGACAACACGGGGCGCCCGTGCGGCACCCGTATAAATCCGGCGCCACA
- 1681 ThrArgGlyValAlaLysAlaValAspPheIleProValGluAsnLeuGluThrThrMet  
GCACCCGTGGAGTGGCTAAGGCGGTGGACTTTATCCCTGTGGAGAACCCTAGAGACAACCA  
CGTGGGCACCTCACCGATTCCGCCACCTGAAATAGGGACACCTCTTGGATCTCTGTTGGT
- 1741 ArgSerProValPheThrAspAsnSerSerProProValValProGlnSerPheGlnVal  
TGAGGTCCCCGGTGTTCACGGATAACTCCTCTCCACCACTAGTGCCCCAGAGCTTCCAGG  
ACTCCAGGGGCCACAAGTGCCTATTGAGGAGAGGTGGTCATCACGGGGTCTCGAAGGTCC
- 1801 AlaHisLeuHisAlaProThrGlySerGlyLysSerThrLysValProAlaAlaTyrAla  
TGGCTCACTCCATGCTCCACAGGCAGCGCAAAGCACCAGGTCCCGGCTGCATATG  
ACCGAGTGGAGGTACGAGGGTGTCCGTGCGCGTTTTCGTGGTTCCAGGGCCGACGTATAC
- 1861 AlaGlnGlyTyrLysValLeuValLeuAsnProSerValAlaAlaThrLeuGlyPheGly  
CAGCTCAGGGCTATAAGGTGCTAGTACTCAACCCCTCTGTTGCTGCAACACTGGGCTTGT  
GTCGAGTCCCGATATTCCACGATCATGAGTTGGGGAGACAACGAGCTTGTGACCCGAAAC
- AlaTyrMetSerLysAlaHisGlyIleAspProAsnIleArgThrGlyValArgThrIle

FIG. 32-2

1921 GTGCTTACATGTCCAAGGCTCATGGGATCGATCCTAACATCAGGACCGGGGTGAGAACA  
 CACGAATGTACAGGTTCCGAGTACCCTAGCTAGGATTGTAGTCCTGGCCCCACTCTTGTT  
 ThrThrGlySerProIleThrTyrSerThrTyrGlyLysPheLeuAlaAspGlyGlyCys  
 1981 TTACCACTGGCAGCCCCATCACGTACTCCACCTACGGCAAGTTCCTTGCCGACGGCGGGT  
 AATGGTGACCGTCGGGGTAGTGCATGAGGTGGATGCCGTTCAAGGAACGGCTGCGCCCA  
 SerGlyGlyAlaTyrAspIleIleIleCysAspGluCysHisSerThrAspAlaThrSer  
 2041 GCTCGGGGGCGCTTATGACATAAATTTGTGACGAGTGCCACTCCACGGATGCCACAT  
 CGAGCCCCCGGAATACTGTATTATTAAACACTGCTACGGTGAGGTGCTACGGTGTA  
 IleLeuGlyIleGlyThrValLeuAspGlnAlaGluThrAlaGlyAlaArgLeuValVal  
 2101 CCATCTTGGGCATCGGCCTGTCTTGACCAAGCAGAGACTGCGGGGGGAGACTGGTTG  
 GGTAGAACCCGTAGCCGTGACAGGAAGTGGTTCGTCTCTGACGCCCCGCTCTGACCAAC  
 LeuAlaThrAlaThrProProGlySerValThrValProHisProAsnIleGluGluVal  
 2161 TGCTCGCCACCGCCACCCCTCCGGGCTCCGTCACTGTGCCCCATCCCAACATCGAGGAGG  
 ACGAGCGGTGGCGGTGGGGAGGCCGAGGCAGTGACACGGGGTAGGGTTGTAGCTCCTCC  
 AlaLeuSerThrThrGlyGluIleProPheTyrGlyLysAlaIleProLeuGluValIle  
 2221 TTGCTCTGTCCACCACCGGAGAGATCCCTTTTTACGGCAAGGCTATCCCCCTCGAAGTAA  
 AACGAGACAGGTGGTGGCCTCTCTAGGGAAAAATGCCGTTCCGATAGGGGGAGCTTCATT  
 LysGlyGlyArgHisLeuIlePheCysHisSerLysLysLysCysAspGluLeuAlaAla  
 2281 TCAAGGGGGGAGACATCTCATCTCTGTTCATTCAAAGAAGAAGTGCGACGAACTCGCG  
 AGTTCCCCCCTCTGTAGAGTAGAAGACAGTAAGTTCTCTTCACGCTGCTTGAGCGGC  
 LysLeuValAlaLeuGlyIleAsnAlaValAlaTyrTyrArgGlyLeuAspValSerVal  
 2341 CAAAGCTGGTGCATTGGGCATCAATGCCGTGGCCTACTACCGCGGTCTTGACGTGTCCG  
 GTTTCGACCAGCGTAACCCGTAGTTACGGCACCGGATGATGGCGCCAGAACTGCACAGGC  
 IleProThrSerGlyAspValValValAlaThrAspAlaLeuMetThrGlyTyrThr  
 2401 TCATCCCGACCAGCGGGATGTGTGTGTGGCAACCGATGCCCTCATGACCGGTATA  
 AGTAGGGCTGGTGGCGCTACAAACAGCAGCACCGTTGGCTACGGGAGTACTGGCGGATAT  
 GlyAspPheAspSerValIleAspCysAsnThrCysValThrGlnThrValAspPheSer  
 2461 CCGGGGACTTCGACTCGGTGATAGACTGCAATACGTGTGTACCCAGACAGTCGATTTCA  
 GCGCGCTGAAGCTGAGCCACTATCTGACGTTATGCACACAGTGGGTCTGTGCTAAAGT  
 LeuAspProThrPheThrIleGluThrIleThrLeuProGlnAspAlaValSerArgThr  
 2521 GCCTTGACCTACCTTCACCATGAGACAATCAGCTCCCCCAGGATGCTGTCTCCCGCA  
 CGGAACCTGGGATGGAAGTGGTAACCTCTGTAGTGCGAGGGGGTCTACGACAGAGGGGT  
 GlnArgArgGlyArgThrGlyArgGlyLysProGlyIleTyrArgPheValAlaProGly  
 2581 CTCAACGTGGGGCAGGACTGGCAGGGGGAAGCCAGGCATCTACAGATTGTGTGGCACCGG  
 GAGTTGCAGCCCGTCTGACCGTCCCCCTCGGTCCGTAGATGTCTAAACACCGTGGCC  
 GluArgProSerGlyMetPheAspSerSerValLeuCysGluCysTyrAspAlaGlyCys  
 2641 GGGAGCGCCCTCCGGCATGTTGACTCGTCCGTCCTCTGTGAGTGCTATGACGCAGGCT  
 CCTCGCGGGGAGGCGGTACAAGCTGAGCAGGCAGGAGACACTACGATACTGCGTCCGA  
 AlaTrpTyrGluLeuThrProAlaGluThrThrValArgLeuArgAlaTyrMetAsnThr  
 2701 GTGCTTGGTATGAGCTCACGCCCGCGAGACTACAGTTAGGCTACGAGCGTACATGAACA  
 CACGAACCACTCGAGTGCGGGCGGCTCTGATGTCAATCOGATGCTGCGATGTACTTGT  
 ProGlyLeuProValCysGlnAspHisLeuGluPheTrpGluGlyValPheThrGlyLeu  
 2761 CCCCGGGGCTTCCCGTGTGCCAGGACCATCTTGAATTTGGGAGGGCGCTTTACAGGCC  
 GGGGCCCCGAAGGGCACACGGTCTCTGGTAGAAGTTAAACCCCTCCCGCAGAAATGTCCGG  
 ThrHisIleAspAlaHisPheLeuSerGlnThrLysGlnSerGlyGluAsnLeuProTyr  
 2821 TCACTCATATAGATGCCACTTCTATCCAGACAAAGCAGAGTGGGGAGAACCTTCTCT  
 AGTGAGTATATCTACGGGTGAAGATAGGGTCTGTTCTGCTCACCCCTCTTGAAGGAA  
 LeuValAlaTyrGlnAlaThrValCysAlaArgAlaGlnAlaProProProSerTrpAsp  
 2881 ACCTGGTAGCGTACCAAGCCACCGTGTGCGCTAGGGCTCAAGCCCCCTCCCCATCGTGGG  
 TGGACCATCGCATGGTTCCGGTGGCACACCGATCCCGAGTTCGGGGAGGGGGTAGCACCC

FIG. 32-3

2941 GlnMetTrpLysCysLeuIleArgLeuLysProThrLeuHisGlyProThrProLeuLeu  
 ACCAGATGTGGAAGTGTGTTGATTGCGCTCAAGCCCACCCTCCATGGGCCAACCCCCCTGC  
 TGGTCTACACCTTCACAACTAAGCGGAGTTCGGGTGGGAGGTACCCGGTTGTGGGGACG

3001 TyrArgLeuGlyAlaValGlnAsnGluIleThrLeuThrHisProValThrLysTyrIle  
 TATACAGACTGGGCGCTGTTTCAAGTAAATCACCTGACGCACCCAGTCACCAAATACA  
 ATATGTCTGACCGCGACAAGTCTTACTTTAGTGGGACTGCGTGGGTTCAGTGGTTTATGT

3061 MetThrCysMetSerAlaAspLeuGluValValThrSerThrTrpValLeuValGlyGly  
 TCATGACATGCATGTGGGCCGACCTGGAGGTGCTCACGAGCACCTGGGTGCTCGTTGGCG  
 AGTACTGTACGTACAGCCGGCTGGACCTCCAGCAGTGTCTGTTGGACCCAGGCAACCGC

3121 ValLeuAlaAlaLeuAlaAlaTyrCysLeuSerThrGlyCysValValIleValGlyArg  
 GCGTCTGGCTGCTTTGGCGCGTATTGCGTGTCAACAGGCTGCGTGGTCATAGTGGGCA  
 CGCAGGACCGACGAAACCGGCGCATAACGGACAGTTGTCCGACGCACCACTATCACCCGT

3181 ValValLeuSerGlyLysProAlaIleIleProAspArgGluValLeuTyrArgGluPhe  
 GGGTGTCTTGTTCGGAAGCCGGCAATCATACCTGACAGGAAGTCTCTACCGAGAGT  
 CCCAGCAGAACAGGCCCTTGGCGGTTAGTATGGACTGTCCCTTCAGGAGATGGCTCTCA

3241 AspGluMetGluGluCysSerGlnHisLeuProTyrIleGluGlnGlyMetMetLeuAla  
 TCGATGAGATGGAAGAGTGTCTCAGCACTTACCGTACATCGAGCAAGGGATGATGCTCG  
 AGCTACTCTACCTTCTCAGGAGTGTGTAATGGCATGTAGTCTGTTCCCTACTACGAGC

3301 GluGlnPheLysGlnLysAlaLeuGlyLeuLeuGlnThrAlaSerArgGlnAlaGluVal  
 CCGAGCAGTTCAAGCAGAAGGCCCTCGGCCCTCTGCAGACCGCGTCCCGTCAGGCAGAGG  
 GGCTCGTCAAGTTCTGTTCCGGGAGCCGGAGGACGTCTGGCGCAGGGCAGTCCGTCTCC

3361 IleAlaProAlaValGlnThrAsnTrpGlnLysLeuGluThrPheTrpAlaLysHisMet  
 TTATCGCCCCCTGTGTCCAGACCACTGGCAAACTCGAGACCTTCTGGGCGAAGCATA  
 AATAGCGGGGACGACAGGTCTGGTTGACCGTTTTTGTAGCTCTGGAAGACCCGCTTCGTAT

3421 TrpAsnPheIleSerGlyIleGlnTyrLeuAlaGlyLeuSerThrLeuProGlyAsnPro  
 TGTGGAACCTTCATCAGTGGGATACAATACTTGGCGGGCTGTGTAACGCTGCCTGGTAACC  
 ACACCTTGAAGTAGTCACCTATGTTATGAACGCCCCGAACAGTTGCGACGGACCATGG

3481 AlaIleAlaSerLeuMetAlaPheThrAlaAlaValThrSerProLeuThrThrSerGln  
 CCGCCATTGCTTCATTGATGGCTTTTACAGCTGCTGTACCAGCCCACTAACCCTAGCC  
 GCGGTAACGAAGTAACCTACGAAAATGTGACGACAGTGGTGGGTGATTGGTGATCGG

3541 ThrLeuLeuPheAsnIleLeuGlyGlyTrpValAlaAlaGlnLeuAlaAlaProGlyAla  
 AAACCTCTCTTCAACATATTGGGGGGGTGGGTGGCTGCCAGCTCGCGGCCCCCGGTG  
 TTTGGGAGGAGAAGTTGTATAACCCCCCACCACCGACGGGTGAGCGGGGGGGGCCAC

3601 AlaThrAlaPheValGlyAlaGlyLeuAlaGlyAlaAlaIleGlySerValGlyLeuGly  
 CCGCTACTGCCTTTGTGGGCGCTGGCTTAGCTGGCGCGCCATCGGCAGTGTGGACTGG  
 GCGATGACGGAAACACCGCGACCGAATCGACCGGGCGGTAGCCGTCAACCTGACC

3661 LysValLeuIleAspIleLeuAlaGlyTyrGlyAlaGlyValAlaGlyAlaLeuValAla  
 GGAAGTCTCATAGACATCTTGCAGGTATGGCGGGGCGTGGCGGGAGCTCTGTGG  
 CCTTCCAGGAGTATCTGTAGGAACGTCCCATACCGCGCCCGCACCGCCCTCGAGAACACC

3721 PheLysIleMetSerGlyGluValProSerThrGluAspLeuValAsnLeuLeuProAla  
 CATTCAAGATCATGAGCGGTGAGGTCCCTCCACGGAGGACCTGGTCAATCTACTGCCCG  
 GTAAGTCTAGTACTGCCACTCCAGGGGAGGTGCTCCTGGACCAGTTAGATGACGGGC

3781 IleLeuSerProGlyAlaLeuValValGlyValValCysAlaAlaIleLeuArgArgHis  
 CCATCTCTCGCCCGGAGCCCTGTAGTCGGCGTGGTCTGTGCAGCAATACTGCGCGCGC  
 GGTAGGAGAGCGGGCCTCGGGAGCATCAGCGCACCAAGACAGTCTTATGACGCGGCGC

3841 ValGlyProGlyGluGlyAlaValGlnTrpMetAsnArgLeuIleAlaPheAlaSerArg  
 ACGTTGGCCCGGCGAGGGGGCAGTGCAGTGGATGAACCGGCTGATAGCTTCGCCTCC  
 TGCAACCGGGCCCGCTCCCCCGTCACTGACCTTGGCCGACTATCGGAAGCGGAGGG

GlyAsnHisValSerProThrHisTyrValProGluSerAspAlaAlaAlaArgValThr

FIG. 32-4

3901 GGGGGAACCATGTTTCCCCACGCACTACGTGCCGGAGAGCGATGCAGCTGCCCCGCGTCA  
 CCCCCTTGTTACAAAGGGGGTGCGTGATGCACGGCCTCTCGCTACGTGACGGGCGCAGT  
 AlaIleLeuSerSerLeuThrValThrGlnLeuLeuArgArgLeuHisGlnTrpIleSer  
 3961 CTGCCATACTCAGCAGCCTCACTGTAACCCAGCTCCTGAGGCGACTGCACCAGTGGATAA  
 GACGGTATGAGTCGTGGAGTGACATTGGGTCGAGGACTCCGCTGACGTGGTCACCTATT  
 SerGluCysThrThrProCysSerGlySerTrpLeuArgAspIleTrpAspTrpIleCys  
 4021 GCTCGGAGTGTAACCACTCCATGCTCGGTTCTCGGTAAGGGACATCTGGGACTGGATAT  
 CGAGCCTCACATGGTGAGGTACGAGGCCAAGGACOGATTCCCTGTAGACCCTGACCTATA  
 GluValLeuSerAspPheLysThrTrpLeuLysAlaLysLeuMetProGlnLeuProGly  
 4081 GCGAGGTGTTGAGCGACTTTAAGACCTGGCTAAAAGCTAAGCTCATGCCACAGCTGCCTG  
 CGCTCCACAACCTCGCTGAAATTCTGGACCGATTTCGATTGAGTACGGTGTGACGGAC  
 IleProPheValSerCysGlnArgGlyTyrLysGlyValTrpArgValAspGlyIleMet  
 4141 GGATCCCTTTGTGTCTCTGCCAGCGCGGGTATAAGGGGGTCTGGCGAGTGGACGGCATCA  
 CCTAGGGGAAACACAGGACGGTGGCGCCATATTCCCCAGACCGCTCACCTGCCGTAGT  
 HisThrArgCysHisCysGlyAlaGluIleThrGlyHisValLysAsnGlyThrMetArg  
 4201 TGCACACTCGCTGCCACTGTGGAGCTGAGATCACTGGACATGTCAAAAACGGGACGATGA  
 ACGTGTGAGCGACGGTGACACCTCGACTCTAGTGACCTGTACAGTTTTGGCCCTGCTACT  
 IleValGlyProArgThrCysArgAsnMetTrpSerGlyThrPheProIleAsnAlaTyr  
 4261 GGATCGTCCGTCTTAGGACCTGCAGGAACATGTGGAGTGGGACCTTCCCCATTAAATGCCT  
 CCTAGCAGCCAGGATCCTGGACGTCTTGTACACCTCACCTGGAAGGGGTAATTACGGA  
 ThrThrGlyProCysThrProLeuProAlaProAsnTyrThrPheAlaLeuTrpArgVal  
 4321 ACACCACGGGCCCTGTACCCCTCTCTGCGCGAAGTACACGTTGCGGCTATGGAGGG  
 TGTGGTGCCCGGGACATGGGGGAAGGACGGGCTTGATGTGAAGCGGATACCTCCC  
 SerAlaGluGluTyrValGluIleArgGlnValGlyAspPheHisTyrValThrGlyMet  
 4381 TGTCTGCAGAGGAATATGTGGAGATAAGGCAGGTGGGGGACTTCCACTACGTGACGGGTA  
 ACAGACGTCTCCTTATACACCTCTATTCCGTCCACCCCTGAAGGTGATGCACTGCCCAT  
 ThrThrAspAsnLeuLysCysProCysGlnValProSerProGluPhePheThrGluLeu  
 4441 TGACTACTGACAATCTCAAATGCCGTGCCAGGTCCCATCGCCCGAATTTTTCACAGAAT  
 ACTGATGACTGTAGAGTTTACGGGCACGGTCCAGGGTAGCGGGCTAAAAAGTGCTTA  
 AspGlyValArgLeuHisArgPheAlaProProCysLysProLeuLeuArgGluGluVal  
 4501 TGGACGGGGTGGCCTACATAGGTTTGCGCCCCCTGCAAGCCCTGCTGCGGGAGGAGG  
 ACCTGCCCAOCCGGATGTATCCAAACGCGGGGGGACGTTGGGGAACGACGCCCTCTCC  
 SerPheArgValGlyLeuHisGluTyrProValGlySerGlnLeuProCysGluProGlu  
 4561 TATCATTCAGAGTAGGACTCCACGAATACCGGTAGGGTCCGAATTACCTTGCAGAGCCG  
 ATAGTAAGTCTCATCTGAGGTGCTTATGGGCCATCCAGCGTTAATGGAACGCTCGGGC  
 ProAspValAlaValLeuThrSerMetLeuThrAspProSerHisIleThrAlaGluAla  
 4621 AACCGGACGTGGCCGTGTGACGTCCATGCTCACTGATCCCTCCCATATAACAGCAGAGG  
 TTGGCCTGCACCGGCACAACCTGCAGGTACGAGTGACTAGGGAGGGTATATTGTCGTCTCC  
 AlaGlyArgArgLeuAlaArgGlySerProProSerValAlaSerSerSerAlaSerGln  
 4681 CGGCCGGGCGAAGGTGGGAGGGGATACCCCTCTGTGGCCAGCTCCTCGGCTAGCC  
 GCGGGCCCGCTTCCAACCGCTCCCTAGTGGGGGAGACACCGGTGAGGAGCCGATCGG  
 LeuSerAlaProSerLeuLysAlaThrCysThrAlaAsnHisAspSerProAspAlaGlu  
 4741 AGCTATCCGCTCCATCTCTCAAGGCAACTTGACCGCTAACCATGACTCCCTGATGCTG  
 TCGATAGGCGAGGTAGAGAGTTCCGTTGAACGTGGCGATTGGTACTGAGGGGACTACGAC  
 LeuIleGluAlaAsnLeuLeuTrpArgGlnGluMetGlyGlyAsnIleThrArgValGlu  
 4801 AGCTCATAGAGCCCAACCTCCTATGGAGGCAGAGATGGGCGGCAACATCACCAGGGTGG  
 TCGAGTATCTCCGTTGGAGGATACCTCCGTCTCTACCGCGGTTGTAGTGGTCCCAAC  
 SerGluAsnLysValValIleLeuAspSerPheAspProLeuValAlaGluGluAspGlu  
 4861 AGTCAGAAAACAAAGTGGTGATTCTGGACTCCTTCGATCCGCTTGTGGCGGAGGAGGACG  
 TCAGTCTTTTTCACCACTAAGACCTGAGGAAGCTAGGCGAACACCGCCTCTCTCTGC

FIG. 32-5

4921 ArgGluIleSerValProAlaGluIleLeuArgLysSerArgArgPheAlaGlnAlaLeu  
 AGCGGGAGATCTCCGTACCCGAGAAATCCTGCGGAAGTCTCGGAGATTGCCCCAGGCCC  
 TCGCCCTCTAGAGGCATGGGCGTCTTTAGGACGCCCTCAGAGCCTCTAAGCGGGTCCGGG  
 4981 ProValTrpAlaArgProAspTyrAsnProProLeuValGluThrTrpLysLysProAsp  
 TGCCCGTTTGGGCGCGGCCGACTATAACCCCCGCTAGTGGAGACGTGGAAAAGCCCCG  
 ACGGGCAAACCCGCGCGGCCCTGATATTGGGGGGCGATCACCTCTGCACCTTTTTCGGGGC  
 5041 TyrGluProProValValHisGlyCysProLeuProProProLysSerProProValPro  
 ACTACGAACCACCTGTGGTCCATGGCTGTCCGCTTCCACCTCCAAAGTCCCTCTCTGTGC  
 TGATGCTTGGTGGACACCAGGTACCGACAGGCGAAGGTGGAGGTTTCAGGGGAGGACACG  
 5101 ProProArgLysLysArgThrValValLeuThrGluSerThrLeuSerThrAlaLeuAla  
 CTCGCCCTCGGAAGAAGCGGACGGTGGTCTCACTGAATCAACCCTATCTACTGCCTTGG  
 GAGGCGGAGCCTTCTTCGCTGCCACCAGGAGTGACTTAGTTGGGATAGATGACGGAACC  
 5161 GluLeuAlaThrArgSerPheGlySerSerSerThrSerGlyIleThrGlyAspAsnThr  
 CCGAGCTCGCCACCAGAAGCTTTGGCAGCTCCTCAACTTCGGGCATTACGGGCGACAATA  
 GGCTCGAGCGGTGGTCTTCGAAACCGTCGAGGAGTTGAAGGCCGTAAATGCCCGCTGTTAT  
 5221 ThrThrSerSerGluProAlaProSerGlyCysProProAspSerAspAlaGluSerTyr  
 CGACAACATCCTCTGAGCCCGCCCTTCTGGCTGCCCCCGGACTCCGACGCTGAGTCCT  
 GCTGTGTAGGAGACTCGGGCGGGGAAGACCACGGGGGGGCTGAGGCTGCGACTCAGGA  
 5281 SerSerMetProProLeuGluGlyGluProGlyAspProAspLeuSerAspGlySerTrp  
 ATTCTCCATGCCCCCTGGAGGGGGAGCCTGGGGATCCGGATCTTAGCGACGGGTCAT  
 TAAGGAGGTACGGGGGGACCTCCCCCTCGGACCCCTAGGCCTAGAATCGCTGCCCAGTA  
 5341 SerThrValSerSerGluAlaAsnAlaGluAspValValCysCysSerMetSerTyrSer  
 GGTCAACGGTCAGTAGTGAGGCCAACGCGGAGGATGTCGTGTGCTGCTCAATGTCTTACT  
 CCAGTTGCCAGTCATCACTCCGGTTGCGCCTCCTACAGCACACGACGAGTTACAGAATGA  
 5401 TrpThrGlyAlaLeuValThrProCysAlaAlaGluGluGlnLysLeuProIleAsnAla  
 CTGGACAGGGCACTCGTCACCCCGTGGCGCGGAAGAACAGAACTGCCCATCAATG  
 GAACCTGTCCGGTGAGCAGTGGGGCACGCGGCGCCTTCTTGTCTTTGACGGGTAGTTAC  
 5461 LeuSerAsnSerLeuLeuArgHisHisAsnLeuValTyrSerThrThrSerArgSerAla  
 CACTAAGCAACTCGTTGCTACGTACCCACAATTGGGTGATTCCACCACCTCACGCAGTG  
 GTGATTGCTTGAGCAACGATGCAGTGGTGTAAACCACATAAGGTGGTGGAGTGCCTCAC  
 5521 CysGlnArgGlnLysLysValThrPheAspArgLeuGlnValLeuAspSerHisTyrGln  
 CTTGCCAAAGGCAGAAGAAAGTCAATTTGACAGACTGCAAGTTCTGGACAGCCATTACC  
 GAACGGTTTTCGTCTTCTTTCAAGTGTAACTGTCTGACGTTCAAGACCTGTCGGTAATGG  
 5581 AspValLeuLysGluValLysAlaAlaAlaSerLysValLysAlaAsnLeuLeuSerVal  
 AGGACGTACTCAAGGAGGTTAAAGCAGCGGCGTCAAAAGTGAAGGCTAACTTGCTATCCG  
 TCCTGCATGAGTTCTCCAATTTCTGTCGCGCAGTTTTCACTTCCGATTGAACGATAGGC  
 5641 GluGluAlaCysSerLeuThrProProHisSerAlaLysSerLysPheGlyTyrGlyAla  
 TAGAGGAAGCTTGACGCTGACGCCCCACACTCAGCCAAATCCAAGTTTGGTTATGGGG  
 ATCTCCTTCGAACGTCGACTGCGGGGTGTGAGTCGGTTTAGGTTCAAACCAATACCCC  
 5701 LysAspValArgCysHisAlaArgLysAlaValThrHisIleAsnSerValTrpLysAsp  
 CAAAAGACGTCCGTTGCCATGCCAGAAAGGCCGTAAACCCACATCAACTCCGTGTGGAAG  
 GTTTTCTGCAGGCAACGGTACGGTCTTTCCGCGCATGGGTGTAGTTGAGGCACACCTTTC  
 5761 LeuLeuGluAspAsnValThrProIleAspThrThrIleMetAlaLysAsnGluValPhe  
 ACCTTCTGGAAGACAATGTAACACCAATAGACACTACCATCATGGCTAAGAACGAGGTTT  
 TGAAGACCTTCTGTTACATTGTGGTTATCTGTGATGGTAGTACCGATTCTTGCTCCAA  
 5821 CysValGlnProGluLysGlyGlyArgLysProAlaArgLeuIleValPheProAspLeu  
 TCTGCGTTTACGCTGAGAAGGGGGGTGTAAGCCAGCTCGTCTCATCGTGTTCGCCGATC  
 AGACGCAAGTCGACTCTTCCCCCAGCATTCGGTCGAGCAGAGTAGCACAAGGGGGCTAG  
 GlyValArgValCysGluLysMetAlaLeuTyrAspValValThrLysLeuProLeuAla

FIG. 32-6



5881 TGGGCGTGC GCGTGTGCGAAAAGATGGCTTTGTACGACGTGGTTACAAAGCTCCCCCTGG  
 ACCCGCACGCGCACACGCTTTTCTACCGAAACATGCTGCACCAATGTTTCGAGGGGAACC  
  
 ValMetGlySerSerTyrGlyPheGlnTyrSerProGlyGlnArgValGluPheLeuVal  
 5941 CCGTGATGGGAAGCTCCTACGGATTCCAATACTCACCAGGACAGCGGGTTGAATTCCTCG  
 GGCCTACCTTCGAGGATGCCTAAGGTTATGAGTGGTCTGTGCGCCAACTTAAGGAGC  
  
 GlnAlaTrpLysSerLysLysThrProMetGlyPheSerTyrAspThrArgCysPheAsp  
 6001 TGCAAGCGTGGAAGTCCAAGAAAACCCCAATGGGGTTCTCGTATGATACCGCTGCTTTG  
 ACGTTCCGACCTTCAGGTTCTTTTGGGGTTACCCCAAGAGCATACTATGGGCGACGAAAC  
  
 SerThrValThrGluSerAspIleArgThrGluGluAlaIleTyrGlnCysCysAspLeu  
 6061 ACTCCACAGTCACTGAGAGCGACATCCGTACGAGGAGGCAATCTACCAATGTTGTGACC  
 TGAGGTGTCAGTGACTCTCGCTGTAGGCATGCCCTCCTCGTTAGATGGTTACAACACTGG  
  
 AspProGlnAlaArgValAlaIleLysSerLeuThrGluArgLeuTyrValGlyGlyPro  
 6121 TCGACCCCAAGCCCGCGTGGCCATCAAGTCCCTCACCAGAGAGGCTTTATGTTGGGGGCC  
 AGCTGGGGGTTCCGGCGCACGGTAGTTTCAGGGAGTGGCTCTCGAAATACAACCCCGG  
  
 LeuThrAsnSerArgGlyGluAsnCysGlyTyrArgArgCysArgAlaSerGlyValLeu  
 6181 CTCTTACCAATTCAAGGGGGGAGAACTGCGGCTATCGCAGGTGCGCGCGAGCGGCGTAC  
 GAGAATGGTTAAGTTCCCCCTCTTGACGCCGATAGCGTCCACGGCGCGCTCGCCGATG  
  
 ThrThrSerCysGlyAsnThrLeuThrCysTyrIleLysAlaArgAlaAlaCysArgAla  
 6241 TGACAACTAGCTGTGGTAACACCCTCACTTGCTACATCAAGGCCGGGCGAGCCTGTGAG  
 ACTGTTGATCGACACCATTGTGGGAGTGAACGATGTAGTTCCGGGCCCGTGGACAGCTC  
  
 AlaGlyLeuGlnAspCysThrMetLeuValCysGlyAspAspLeuValValIleCysGlu  
 6301 CCGCAGGGCTCCAGGACTGCACCATGCTCGTGTGTGGCGACGACTTAGTCGTTATCTGTG  
 GCGTCCCGAGGTCTGACGTGGTACGAGCACACACCGCTGCTGAATCAGCAATAGACAC  
  
 SerAlaGlyValGlnGluAspAlaAlaSerLeuArgAlaPheThrGluAlaMetThrArg  
 6361 AAAGCGCGGGGGTCCAGGAGGACGCGCGAGCCTGAGAGCCTTCACGGAGGCTATGACCA  
 TTTCCGCGCCCCAGGTCTCTCGCGCGCTCGGACTCTCGGAAGTGCTCCGATACTGGT  
  
 TyrSerAlaProProGlyAspProProGlnProGluTyrAspLeuGluLeuIleThrSer  
 6421 GGTACTCCGCCCCCTGGGGACCCCAACAGGAGTACGACTTGGAGCTCATACAT  
 CCATGAGCGGGGGGACCCTGGGGGGTGTGGTCTTATGCTGAACCTGAGTATTGTA  
  
 CysSerSerAsnValSerValAlaHisAspGlyAlaGlyLysArgValTyrTyrLeuThr  
 6481 CATGCTCTCCAACGTGTGAGTGGCCACGACGGCGCTGGAAAGAGGGTCTACTACCTCA  
 GTACGAGGAGGTTGCACAGTCAGCGGGTGTGCGCGACCTTTCTCCAGATGATGGAGT  
  
 ArgAspProThrThrProLeuAlaArgAlaAlaTrpGluThrAlaArgHisThrProVal  
 6541 CCGTGACCTACAACCCCCCTCGCGAGAGCTGCGTGGGAGACAGCAAGACACACTCCAG  
 GGGCACTGGGATGTTGGGGGGAGCGCTCTCGACGACCTCTGTGCTTCTGTGTGAGGTC  
  
 AsnSerTrpLeuGlyAsnIleIleMetPheAlaProThrLeuTrpAlaArgMetIleLeu  
 6601 TCAATTCCTGGCTAGGCAACATAATCATGTTTGGCCCCACACTGTGGGCGAGGATGATAC  
 AGTTAAGGACCGATCCGTTGTATTAGTACAAACGGGGGTGTGACACCGCTCCTACTATG  
  
 MetThrHisPhePheSerValLeuIleAlaArgAspGlnLeuGluGlnAlaLeuAspCys  
 6661 TGATGACCATTTCTTTAGCGTCTTATAGCCAGGACAGCTTGAACAGGCCCTCGATT  
 ACTACTGGGTAAAGAAATCGCAGGAATATCGGTCCTGGTGAACCTGTGCGGGAGCTAA  
  
 GluIleTyrGlyAlaCysTyrSerIleGluProLeuAspLeuProProIleIleGlnArg  
 6721 GCGAGATCTACGGGGCCTGCTACTCCATAGAACCCTTGATCTACCTCCAATCATTCAA  
 CGCTCTAGATGCCCCGGACGATGAGGTATCTTGGTGAACCTAGATGGAGGTTAGTAAGTTT  
  
 Leu  
 6781 GACTC  
 CTGAG

FIG. 32-7

FIG. 47-1 COMBINED ORF OF DNAs K9-1 through 15e

GlyCysProGluArgLeuAlaSerCysArgProLeuThrAspPheAspGlnGlyTrpGly  
 1 CAGGCTGCCTGAGAGGCTAGCCAGCTGCCAGCCCTTACCGATTTTGACCAGGGCTGGG  
 GTCCGACAGGACTCTCCGATCGGTCGACGGCTGGGGAATGGCTAAACTGGTCCCCAGCC

ProIleSerTyrAlaAsnGlySerGlyProAspGlnArgProTyrCysTrpHisTyrPro  
 61 GCCCTATCAGTTATGCCAACGGAAGCGCCCCGACCAGCGCCCTACTGCTGGCACTACC  
 CGGGATAGTCAATACGGTTGCCTTCGCCGGGGCTGGTCGCGGGGATGACGACCGTGATGG

ProLysProCysGlyIleValProAlaLysSerValCysGlyProValTyrCysPheThr  
 121 CCCCCAAACCTTGCGGTATTGTGCCCGGAAGAGTGTGTGTGGTCCGGTATATTGCTTCA  
 GGGGTTTTTGAACGCCATAACACGGGCGCTTCTCACACACACCAGGCCATATAACGAAGT

ProSerProValValValGlyThrThrAspArgSerGlyAlaProThrTyrSerTrpGly  
 181 CTCCCAGCCCCGTGGTGGTGGGAACGACCGACAGGTGCGGGCGCGCCACCTACAGCTGGG  
 GAGGGTCGCGGGCACCACCACCCTTGCTGGCTGTCCAGCCCGCGCGGGTGGATGTGACCC

GluAsnAspThrAspValPheValLeuAsnAsnThrArgProProLeuGlyAsnTrpPhe  
 241 GTGAAATGATACGGACGTCTTCGTCTTAACAATACCAGGCCACCGTGGGCAATTGGT  
 CACTTTTACTATGCTGCAGAAGCAGGAATTGTTATGGTCCGGTGGCGACCCGTTAACCA

GlyCysThrTrpMetAsnSerThrGlyPheThrLysValCysGlyAlaProProCysVal  
 301 TCGGTTGTACCTGGATGAACCTCACTGGATTACCAAAGTGTGCGGAGCGCCTCCTGTG  
 AGCCAACATGGACCTACTTGAGTTGACCTAAGTGGTTTCACACGCCTCGCGGAGGAACAC

IleGlyGlyAlaGlyAsnAsnThrLeuHisCysProThrAspCysPheArgLysHisPro  
 361 TCATCGGAGGGGCGGCAACAACCCCTGCACCTGCCCCACTGATTGCTTCCGCAAGCATC  
 AGTAGCCTCCCCGCCGTTGTTGTGGGACGTGACGGGGTACTAACGAAGGCGTTCGTAG

AspAlaThrTyrSerArgCysGlySerGlyProTrpIleThrProArgCysLeuValAsp  
 421 CGGACGCCACATACTCTCGGTGCGGCTCCGGTCCCTGGATCACACCCAGGTGCCTGGTGC  
 GCCTGCGGTGTATGAGAGCCACGCCGAGGCCAGGACCTAGTGTGGGTCCACGGACCAGC

TyrProTyrArgLeuTrpHisTyrProCysThrIleAsnTyrThrIlePheLysIleArg  
 481 ACTACCCGTATAGGCTTTGGCATTATCCTTGTACCATCAACTACCCATATTTAAATCA  
 TGATGGGCATATCCGAAACCGTAATAGGAACATGGTAGTTGATGTGGTATAAATTTAGT

MetTyrValGlyGlyValGluHisArgLeuGluAlaAlaCysAsnTrpThrArgGlyGlu  
 541 GGATGTACGTGGGAGGGGTGGAACACAGGCTGGAAGCTGCCTGCAACTGGACGCGGGGCG  
 CCTACATGCACCCTCCCCAGCTTGTGTCCGACCTTCGACGGACGTTGACCTGCGCCCCGC

ArgCysAspLeuGluAspArgAspArgSerGluLeuSerProLeuLeuLeuThrThrThr  
 601 AACGTTGCGATCTGGAAGACAGGGACAGGTCCGAGCTCAGCCCGTTACTGCTGACCACTA  
 TTGCAACGCTAGACCTTCTGTCCCTGTCCAGGCTCGAGTCGGGCAATGACGACTGGTGAT

GlnTrpGlnValLeuProCysSerPheThrThrLeuProAlaLeuSerThrGlyLeuIle  
 661 CACAGTGGCAGGTCTCCCGTGTTCCTTACAACCTTACCAGCCTTGTCCACCGGCCTCA  
 GTGTCAACGCTCAGGAGGGCACAAGGAAGTGTGGGATGGTGGGAACAGGTGGCCGGAGT

HisLeuHisGlnAsnIleValAspValGlnTyrLeuTyrGlyValGlySerSerIleAla  
 721 TCCACCTCCACCAGAACATTGTGGACGTGCAGTACTTGTACGGGGTGGGGTCAAGCATCG  
 AGGTGGAGGTGGTCTTGTAAACCTGCACGTCAATGAACATGCCCCACCCAGTTCGTAGC

SerTrpAlaIleLysTrpGluTyrValValLeuLeuPheLeuLeuAlaAspAlaArg  
 781 CGTCCTGGGCCATTAGTGGGAGTACGTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT  
 GCAGGACCCGGTAATTCACCTCATGTCAGCAAGAGGACAAGGAAGACGAACGCTCTGCGCG

ValCysSerCysLeuTrpMetMetLeuLeuIleSerGlnAlaGluAlaAlaLeuGluAsn  
 841 GCGTCTGCTCCTGCTTGTGGATGATGCTACTCATATCCCAAGCGGAGGCGGCTTTGGAGA  
 CGCAGACGAGGACGAACACCTACTACGATGAGTATAGGGTTCGCTCCGCCGAAACCTCT

LeuValIleLeuAsnAlaAlaSerLeuAlaGlyThrHisGlyLeuValSerPheLeuVal  
 901 ACCTCGTAATACTTAATGCAGCATCCCTGGCCGGGACGCACGGTCTTGTATCCTTCCTCG  
 TGGAGCATTATGAATTACGTCGTAGGGACCGGCCCTGCGTGCCAGAACATAGGAAGGAGC

- 961 PhePheCysPheAlaTrpTyrLeuLysGlyLysTrpValProGlyAlaValTyrThrPhe  
TGTTCTTCTGCTTTGCATGGTATTTGAAGGGTAAGTGGGTGCCCCGAGCGGTCTACACCT  
ACAAGAAGACGAAACGTACCATAAACTTCCCATTCACCCACGGGCCCTCGCCAGATGTGGA
- 1021 TyrGlyMetTrpProLeuLeuLeuLeuLeuAlaLeuProGlnArgAlaTyrAlaLeu  
TCTACGGGATGTGGCCTCTCCTGCTCCTGTGGCGTTGCCCCAGCGGGCGTACGGC  
AGATGCCCTACACCGGAGAGGAGGACGAGGACAACCGCAACGGGGTTCGCCCATGCGCG
- 1081 AspThrGluValAlaAlaSerCysGlyGlyValValLeuValGlyLeuMetAlaLeuThr  
TGGACACGGAGGTGGCCGCGTGTGGCGGTGTGTCTCGTGGGTGATGGCGCTGA  
ACCTGTGCCTCCACCGGCGCAGCACACCGCCACAACAAGAGCAGCCAACTACCGCGACT
- 1141 LeuSerProTyrTyrLysArgTyrIleSerTrpCysLeuTrpTrpLeuGlnTyrPheLeu  
CTCTGTCACCATATTACAAGCGCTATATCAGCTGGTGCTGTGGTGGCTTCAGTATTTTC  
GAGACAGTGGTATAATGTTCCGATATAGTCGACCACGAACACCACOGAAGTCATAAAAG
- 1201 ThrArgValGluAlaGlnLeuHisValTrpIleProProLeuAsnValArgGlyGlyArg  
TGACCAGAGTGAAGCGCAACTGCACGTGTGGATTCCCCCCTCAACGTCCGAGGGGGC  
ACTGGTCTCACCTTCGGGTGACGTGCACACCTAAGGGGGGAGTTGCAGGCTCCCCCG
- 1261 AspAlaValIleLeuLeuMetCysAlaValHisProThrLeuValPheAspIleThrLys  
GCGACGCGTCATCTTACTCATGTGTGTGTACACCGACTCTGGTATTTGACATCACCA  
CGTGCAGCAGTAGAATGAGTACACACGACATGTGGGCTGAGACCATAAACTGTAGTGGT
- 1321 LeuLeuLeuAlaValPheGlyProLeuTrpIleLeuGlnAlaSerLeuLeuLysValPro  
AATTGCTGTGGCGCTCTTCGGACCCCTTTGGATTCTTCAAGCCAGTTTGCTTAAAGTAC  
TTAACGACGACCGGCAGAACCTGGGGAACCTAAGAAGTTCGGTCAAACGAATTTTCATG
- 1381 TyrPheValArgValGlnGlyLeuLeuArgPheCysAlaLeuAlaArgLysMetIleGly  
CCTACTTTGTGGCGCTCCAAGGCCTTCTCCGTTCTGCGCGTTAGCGCGGAAGATGATCG  
GGATGAAACACGCGCAGGTTCCGGAAGAGGCCAAGACGCGCAATCGCGCCTTCTACTAGC
- 1441 GlyHisTyrValGlnMetValIleIleLysLeuGlyAlaLeuThrGlyThrTyrValTyr  
GAGGCCATTACGTGCAAATGGTCATCATTAAAGTTAGGGGCGCTTACTGGCACCTATGTTT  
CTCCGGTAATGCACGTTTACCAGTAGTAATTCAATCCCGGAATGACCGTGGATACAA
- 1501 AsnHisLeuThrProLeuArgAspTrpAlaHisAsnGlyLeuArgAspLeuAlaValAla  
ATAACCATCTCACTCCTCTTCGGGACTGGGCGCACACCGCTTGGCAGATCTGGCCGTGG  
TATTGGTAGAGTGAGGAGAAGCCCTGACCCGCGTGTGCGGAACGCTCTAGACCGGCACC
- 1561 ValGluProValValPheSerGlnMetGluThrLysLeuIleThrTrpGlyAlaAspThr  
CTGTAGAGCCAGTCGTCTTCTCCCAAATGGAGACCAAGCTCATCACGTGGGGGGCAGATA  
GACATCTCGGTACGAGAGAGGGTTTACCTCTGGTTCGAGTAGTGACCCCCCGTCTAT
- 1621 AlaAlaCysGlyAspIleIleAsnGlyLeuProValSerAlaArgArgGlyArgGluIle  
CCGCGCGTGGGTGACATCATCAACGGCTTGCCGTGTTCCGCCCCGAGGGGCCGGGAGA  
GGCGGCGCACGCCACTGTAGTAGTTGCCGAACGGACAAAGGGGGCGTCCCCGGCCCTCT
- 1681 LeuLeuGlyProAlaAspGlyMetValSerLysGlyTrpArgLeuLeuAlaProIleThr  
TACTGCTGGGGCCAGCCGATGGAATGGTCTCCAAGGGGTGGAGGTGCTGGCGCCCATCA  
ATGACGAGCCCGGTCCGCTACCTTACCAGAGGTCCCCACCTCCAACGACCGCGGGTAGT
- 1741 AlaTyrAlaGlnGlnThrArgGlyLeuLeuGlyCysIleIleThrSerLeuThrGlyArg  
CGGCGTACGCCAGCAGACAAGGGGCTCCTAGGGTGCATAATCACCAGCCTAAGTGGCC  
GCCGATGCGGGTCTGTCTTCCCCGGAGGATCCACGTATTAGTGGTGGATTGACCGG
- 1801 AspLysAsnGlnValGluGlyGluValGlnIleValSerThrAlaAlaGlnThrPheLeu  
GGGACAAAAACCAAGTGGAGGTGAGGTCCAGATTGTGTAAGTCTGCCAAACCTTCC  
CCCTGTTTTTGGTTACCTCCCACTCCAGGTCTAACACAGTTGACGACGGGTTTGGAAAG
- 1861 AlaThrCysIleAsnGlyValCysTrpThrValTyrHisGlyAlaGlyThrArgThrIle  
TGGCAACGTGCATCAATGGGGTGTGCTGGACTGTCTACCAAGGGGCGGAACGAGGACCA  
ACCGTTGCACGTAGTTACCCACACGACCTGACAGATGGTCCCCGGCCTTGCTCCTGGT
- AlaSerProLysGlyProValIleGlnMetTyrThrAsnValAspGlnAspLeuValGly

FIG. 47-2

- 1921 TCGCGTCACCCAAGGGTCTGTATCCAGATGTATACCAATGTAGACCAAGACCTTGTGG  
AGCGCAGTGGGTTCACAGGACAGTAGGTCTACATATGGTTACATCTGGTTCTGGAACACC
- 1981 TrpProAlaProGlnGlySerArgSerLeuThrProCysThrCysGlySerSerAspLeu  
GCTGGCCCGCTCCGCAAGGTAGCCGCTCATTGACACCTGCACCTGCGGCTCCTCGGACC  
CGACCGGGCGAGGCGTTCATCGGCGAGTAACTGTGGGACGTGAACGCCGAGGAGCCTGG
- 2041 TyrLeuValThrArgHisAlaAspValIleProValArgArgArgGlyAspSerArgGly  
TTTACCTGGTCACGAGGCACGCCGATGTCTATTCCTGCGCCGCGGGGTGATAGCAGGG  
AAATGGACCAAGTGTCTCGTGGGCTACAGTAAGGGCACGCGGCCGCCCACTATCGTCCC
- 2101 SerLeuLeuSerProArgProIleSerTyrLeuLysGlySerSerGlyGlyProLeuLeu  
GCAGCCTGTGTGCGCCCGGCCCATTTCTTACTTGAAAGGCTCCTCGGGGGGTTCGCTGT  
CGTCGGACGACAGCGGGCGGGGTAAAGGATGAACCTTCCGAGGAGCCCCCAGGCGACA
- 2161 CysProAlaGlyHisAlaValGlyIlePheArgAlaAlaValCysThrArgGlyValAla  
TGTGCCCCGCGGGGCACCGCTGGGCATATTTAGGGCCGCGGTGTGCACCCGTGGAGTGG  
ACACGGGGCGCCCCGTGCGGCACCCGTATAAATCCCGGCCACACGTGGGCACCTCACC
- 2221 LysAlaValAspPheIleProValGluAsnLeuGluThrThrMetArgSerProValPhe  
CTAAGGCGGTGGACTTTATCCCTGTGGAGAACCTAGAGACAACCATGAGGTCCCCGGTGT  
GATTCGCCACCTGAAATAGGGACACCTCTTGATCTCTGTTGTTACTCCAGGGGCCACA
- 2281 ThrAspAsnSerSerProProValValProGlnSerPheGlnValAlaHisLeuHisAla  
TCACGGATAACTCCTCTCCACCAGTAGTGCCCCAGAGCTTCCAGGTGGCTCACCTCCATG  
AGTGCTATTGAGGAGAGGTGGTTCATCACGGGTCTCGAAGGTCCACCGAGTGGAGGTAC
- 2341 ProThrGlySerGlyLysSerThrLysValProAlaAlaTyrAlaAlaGlnGlyTyrLys  
CTCCACAGGCAGCGGCAAAAGCACCAAGGTCCCGGCTGCATATGCAGCTCAGGGCTATA  
GAGGGTGTCCGTGCGCGTTTTCGTGGTTCCAGGGCCGACGTATACGTGAGTCCCGATAT
- 2401 ValLeuValLeuAsnProSerValAlaAlaThrLeuGlyPheGlyAlaTyrMetSerLys  
AGGTGCTAGTACTCAACCCTCTGTTGTGCAACACTGGGCTTGGTGCTTACATGTCCA  
TCCACGATCATGAGTTGGGAGACAACGACGTTGTGACCCGAAACCAATGTACAGGT
- 2461 AlaHisGlyIleAspProAsnIleArgThrGlyValArgThrIleThrThrGlySerPro  
AGGCTCATGGGATCGATCTTAACATCAGGACCGGGGTGAGAACAATTACCACTGGCAGCC  
TCCGAGTACCCTAGCTAGGATTGTAGTCTGGCCCCACTCTTGTTAATGGTGACCGTCGG
- 2521 IleThrTyrSerThrTyrGlyLysPheLeuAlaAspGlyGlyCysSerGlyGlyAlaTyr  
CCATCAGTACTCCACTACGGCAAGTTCCTTGCCGACGGCGGGTGCTCGGGGGGCGCTT  
GGTAGTGCATGAGGTGGATGCCGTTCAAGGAACGGCTGCGCCACGAGCCCCCGCGAA
- 2581 AspIleIleIleCysAspGluCysHisSerThrAspAlaThrSerIleLeuGlyIleGly  
ATGACATAATAATTTGTGACGAGTGCCACTCCAAGGATGCCACATCCATCTTGGGCATCG  
TACTGTATTATTAAACACTGCTCACGGTGAGGTGCTACGGTGTAGGTAGAACCCTAGC
- 2641 ThrValLeuAspGlnAlaGluThrAlaGlyAlaArgLeuValValLeuAlaThrAlaThr  
GCACTGTCTTGACCAAGCAGAGACTGCGGGGCGAGACTGGTTGTGCTCGCCACCGCCA  
CGTGACAGGAAGTGGTTCTCTGACGCCCCGCTCTGACCAACACGAGCGGTGGCGGT
- 2701 ProProGlySerValThrValProHisProAsnIleGluGluValAlaLeuSerThrThr  
CCCCTCGGGCTCCGTCACGTGCCCCATCCCAACATCGAGGAGGTGCTCTGTCCACCA  
GGGGAGGCCCGAGGCAGTGACACGGGGTAGGGTTGTAGCTCCTCCAACGAGACAGGTGGT
- 2761 GlyGluIleProPheTyrGlyLysAlaIleProLeuGluValIleLysGlyGlyArgHis  
COGGAGAGATCCCTTTTTACGGCAAGGCTATCCCTCGAAGTAATCAAGGGGGGAGAC  
GGCCTCTCTAGGGAATAATGCCGTTCCGATAGGGGGAGCTTCATTAGTTCCCCCCTCTG
- 2821 LeuIlePheCysHisSerLysLysLysCysAspGluLeuAlaAlaLysLeuValAlaLeu  
ATCTCATCTTCTGTATTCAAAGAAGAAGTGCAGCAACTCGCCGCAAAGCTGGTTCGCAT  
TAGAGTAGAAGACAGTAAGTTTCTTCTTACGCTGCTTGAGCGGCGTTTCGACACAGCGTA
- 2881 GlyIleAsnAlaValAlaTyrTyrArgGlyLeuAspValSerValIleProThrSerGly  
TGGGCATCAATGCCGTGGCCTACTACCGGCTCTGACGTGTCCGTATCCCGACCAGCG  
ACCGTAGTTACGGCACCGGATGATGGCGCCAGAACTGCACAGGCAGTAGGGCTGGTCCG

FIG. 47-3

- 2941 AspValValValValAlaThrAspAlaLeuMetThrGlyTyrThrGlyAspPheAspSer  
GCGATGTTGTCGTCGTGGCAACCGATGCCCTCATGACCGGTATACCGGCGACTTCGACT  
GGCTACAACAGCAGCACCGTTGGCTACGGGAGTACTGGCCGATATGGCCGCTGAAGCTGA
- 3001 ValIleAspCysAsnThrCysValThrGlnThrValAspPheSerLeuAspProThrPhe  
CGGTGATAGACTGCAATACGTGTGTACCCAGACAGTCGATTTCAGCCTTGACCCTACCT  
GCCACTATCTGACGTTATGCACACAGTGGGTCTGTCTAGCTAAAGTCGGAAGTGGGATGGA
- 3061 ThrIleGluThrIleThrLeuProGlnAspAlaValSerArgThrGlnArgArgGlyArg  
TCACCATTTAGACAATCAGCTCCCCAGGATGCTGTCTCCCGCACTCAACGTCGGGGCA  
AGTGGTAACTCTGTTAGTGGAGGGGGTCTACGACAGAGGGCGTGAGTTGCAGCCCCGT
- 3121 ThrGlyArgGlyLysProGlyIleTyrArgPheValAlaProGlyGluArgProSerGly  
GGACTGGCAGGGGAAGCCAGGCATCTACAGATTGTGTGGCACCGGGGAGCGCCCCCTCCG  
CCTGACCGTCCCCCTTCGGTCCGTAGATGTCTAAACACCGTGGCCCCCTCGCGGGGAGGC
- 3181 MetPheAspSerSerValLeuCysGluCysTyrAspAlaGlyCysAlaTrpTyrGluLeu  
GCATGTTGACTCGTCCGTCTCTGTGAGTGCTATGACGCAGGCTGTCTTGGTATGAGC  
CGTACAAGCTGAGCAGGCAGGAGACACTACGATACTGCGTCCGACACGAACCATACTCG
- 3241 ThrProAlaGluThrThrValArgLeuArgAlaTyrMetAsnThrProGlyLeuProVal  
TCACGCCCCGCGAGACTACAGTTAGGCTACGAGCGTACATGAACCCCCGGGGCTTCCCG  
AGTGGGGCGGCTCTGATGTCAATCCGATGCTCGCATGTACTTGTGGGGCCCCGAAGGGC
- 3301 CysGlnAspHisLeuGluPheTrpGluGlyValPheThrGlyLeuThrHisIleAspAla  
TGTGCCAGGACCATCTTGAATTTGGGAGGGCGTCTTTACAGGCCTCACTCATATAGATG  
ACACGCTCTGGTAGAACTTAAACCTCCCGCAGAAATGTCCGAGTGAGTATATCTAC
- 3361 HisPheLeuSerGlnThrLysGlnSerGlyGluAsnLeuProTyrLeuValAlaTyrGln  
CCCACCTTCTATCCAGACAAAGCAGAGTGGGGAGAACCTTCCTTACCTGGTAGCGTACC  
GGGTGAAAGATAGGGTCTGTTTCGTCTCACCCCTCTTGAAGGAATGGACCATCGCATGG
- 3421 AlaThrValCysAlaArgAlaGlnAlaProProProSerTrpAspGlnMetTrpLysCys  
AAGCCACCGTGTGCGCTAGGGCTCAAGCCCTCCCCCATCGTGGGACCAGATGTGGAAGT  
TTCGGTGGCACACGCGATCCCGAGTTCGGGGAGGGGTAGCACCCCTGGTCTACACCTTCA
- 3481 LeuIleArgLeuLysProThrLeuHisGlyProThrProLeuLeuTyrArgLeuGlyAla  
GTTTGATTCGCTCAAGCCACCTCCATGGGCCAACCCCTGCTATACAGACTGGGGC  
CAAACCTAAGCGGAGTTCGGGTGGGAGGTACCCGGTTGTGGGGACGATATGTCTGACCCGC
- 3541 ValGlnAsnGluIleThrLeuThrHisProValThrLysTyrIleMetThrCysMetSer  
CTGTTCAAGATGAAATCACCTGACGCACCCAGTCACCAAATACATCATGACATGCATGT  
GACAAGTCTTACTTTAGTGGGACTGCGTGGGTCAGTGGTTTATGTAGTACTGTACGTACA
- 3601 AlaAspLeuGluValValThrSerThrTrpValLeuValGlyGlyValLeuAlaAlaLeu  
CGGCCGACCTGGAGGTCGTACGAGCACCTGGGTGCTCGTTGGCGGCGTCTGGCTGCTT  
GCCGGCTGGACCTCCAGCAGTGTCTGTGGACCCACGAGCAACCGCCGAGGACCGACGAA
- 3661 AlaAlaTyrCysLeuSerThrGlyCysValValIleValGlyArgValValLeuSerGly  
TGGCCGCTATTGCTGTCAACAGGCTGCGTGGTCATAGTGGGCAGGGTCTGCTTGTCCG  
ACCGGCCCATACGGACAGTTGTCCGACGCACAGTATACCCGTCCAGCAGAACAGGC
- 3721 LysProAlaIleIleProAspArgGluValLeuTyrArgGluPheAspGluMetGluGlu  
GGAAGCCGGCAATCATACCTGACAGGGAAGTCTCTACCGAGAGTTCGATGAGATGGAAG  
CCTTCGGCCGTTAGTATGGACTGTCCCTCAGGAGATGGCTCTCAAGCTACTCTACCTTC
- 3781 CysSerGlnHisLeuProTyrIleGluGlnGlyMetMetLeuAlaGluGlnPheLysGln  
AGTGCTCTCAGCACTTACCGTACATCGAGCAAGGGATGATGCTCGCCGAGCAGTTCAAGC  
TCACGAGAGTCTGTAATGGCATGTAGCTCGTTCCTACTACGAGCGGCTCGTCAAGTTCCG
- 3841 LysAlaLeuGlyLeuLeuGlnThrAlaSerArgGlnAlaGluValIleAlaProAlaVal  
AGAAGGCCCTCGGCCTCTGCGAGACCGGCTCCCGTCAGGCAGAGGTTATCGCCCCCTGCTG  
TCTTCCGGGAGCGGAGGACGTCTGGCGCAGGGCAGTCTGCTCCAAAGCGGGGACGAC
- GlnThrAsnTrpGlnLysLeuGluThrPheTrpAlaLysHisMetTrpAsnPheIleSer

FIG. 47-4

- 3901 TCCAGACCAACTGGCAAACTCGAGACCTTCTGGGCGAAGCATATGTGGAACCTTCATCA  
AGGCTCGGTGACCGTTTTTGAGCTCTGGAAGACCGCTTCGTATACACCTTGAAGTAGT
- GlyIleGlnTyrLeuAlaGlyLeuSerThrLeuProGlyAsnProAlaIleAlaSerLeu  
3961 GTGGGATACAATACTTGGCGGGCTGTCAACGCTGCCTGGTAACCCGCCATTGCTTCAT  
CACCTATGTTATGAACCGCCGAACAGTTGCGACGGACCATTTGGGGCGTAACGAAGTA
- MetAlaPheThrAlaAlaValThrSerProLeuThrThrSerGlnThrLeuLeuPheAsn  
4021 TGATGGCTTTTACAGCTGCTGTACCAGCCACTAACCCTAGCCAAACCTCTCTCA  
ACTACCGAAAATGTCGACGACAGTGGTCGGGTGATTGGTGATCGGTTTGGGAGGAGAAGT
- IleLeuGlyGlyTrpValAlaAlaGlnLeuAlaAlaProGlyAlaAlaThrAlaPheVal  
4081 ACATATTGGGGGGTGGGTGGCTGCCCAGCTCGCCGCCCCGGTGCCGCTACTGCCTTTG  
TGATAACCCCCCACCACCGACGGGTCGAGCGCGGGGGCCACGGCGATGACGGAAC
- GlyAlaGlyLeuAlaGlyAlaAlaIleGlySerValGlyLeuGlyLysValLeuIleAsp  
4141 TGGGCGCTGGCTTAGCTGGCGCCGCGCATCGGCAGTGTGGACTGGGGAAGGTCTTCATAG  
ACCCGCGACCGAATCGACCGCGCGGTAGCCGTCAACCTGACCCCTTCCAGGAGTATC
- IleLeuAlaGlyTyrGlyAlaGlyValAlaGlyAlaLeuValAlaPheLysIleMetSer  
4201 ACATCCTTGCAGGGTATGCGCGGGCGTGGCGGGAGCTCTTGTGGCATTCAAGATCATGA  
TGTAGGAACGTCCCATACCGCGCCCGACCGCCCTCGAGAACACCGTAAGTTCTAGTACT
- GlyGluValProSerThrGluAspLeuValAsnLeuLeuProAlaIleLeuSerProGly  
4261 GCGGTGAGGTCCCTCCACGGAGGACCTGGTCAATCTACTGCCCGCCATCCTCTCGCCCG  
CGCCACTCCAGGGGAGGTGCTCTGGACAGTTAGATGACGGGCGGTAGGAGAGCGGGC
- AlaLeuValValGlyValValCysAlaAlaIleLeuArgArgHisValGlyProGlyGlu  
4321 GAGCCCTCGTAGTCGGCGTGGTCTGTGCAGCAATACTGCGCCGGCACGTTGGCCCGGGCG  
CTCGGGAGCATCAGCCGCACCAGACACGTCGTTATGACGCGCGCGTCAACCGGGCCCGC
- GlyAlaValGlnTrpMetAsnArgLeuIleAlaPheAlaSerArgGlyAsnHisValSer  
4381 AGGGGGCAGTGCAGTGGATGAACCGGCTGATAGCCTTCGCCTCCCGGGGAACCATGTTT  
TCCCCCGTCACGTCACCTACTTGGCCGACTATCGGAAGCGGAGGGCCCCCTTGGTACAAA
- ProThrHisTyrValProGluSerAspAlaAlaAlaArgValThrAlaIleLeuSerSer  
4441 CCCCCACGCACTACGTGCCGGAGAGCGATGCAGCTGCCCGCGTCACTGCCATACTCAGCA  
GGGGGTGCGTGATGCACGGCCTCTCGCTACGTGCAGGGGCGCAGTGACGGTATGAGTCGT
- LeuThrValThrGlnLeuLeuArgArgLeuHisGlnTrpIleSerSerGluCysThrThr  
4501 GCCTCACTGTAACCCAGCTCCTGAGGCGACTGCACCACTGGATAAGCTCGGAGTGATACCA  
CGGAGTGACATTGGGTGAGGACTCGCTGACGTGGTCACCTATTCGAGCCTCACATGGT
- ProCysSerGlySerTrpLeuArgAspIleTrpAspTrpIleCysGluValLeuSerAsp  
4561 CTCCATGCTCCGGTTCTGGCTAAGGGACATCTGGGACTGGATATGCGAGGTGTTGAGCG  
GAGGTACGAGGCCAAGGACCGATTCCCTGTAGACCCTGACCTATACGCTCCACAACCTCGC
- PheLysThrTrpLeuLysAlaLysLeuMetProGlnLeuProGlyIleProPheValSer  
4621 ACTTTAAGACCTGGCTAAAAGCTAAGCTCATGCCACAGCTGCCTGGGATCCCCCTTTGTGT  
TGAAATTCTGGACCGATTTCGATTTCGAGTACGGTGTCGACGGACCTAGGGGAAACACA
- CysGlnArgGlyTyrLysGlyValTrpArgValAspGlyIleMetHisThrArgCysHis  
4681 CCTGCCAGCGCGGTATAAGGGGGTCTGGCGAGTGGACGGCATCATGCACACTCGCTGCC  
GGACGGTCCGCCCATTATCCCCCAGACCGCTCACCTGCCGTAGTACGTGTGAGCGACGG
- CysGlyAlaGluIleThrGlyHisValLysAsnGlyThrMetArgIleValGlyProArg  
4741 ACTGTGGAGCTGAGATCACTGGACATGTCAAAAACGGGACGATGAGGATCGTCGGTCTTA  
TGACACCTCGACTCTAGTGACCTGTACAGTTTTTGGCCCTGCTACTCTAGCAGCCAGGAT
- ThrCysArgAsnMetTrpSerGlyThrPheProIleAsnAlaTyrThrThrGlyProCys  
4801 GGACCTGCAGGAACATGTGGAGTGGGACCTTCCCCATTATGCCTACACCACGGGCCCCCT  
CCTGGACGTCCTGTACACCTACCCCTGGAAGGGGTAATTACGGATGTGGTGCCCGGGGA
- ThrProLeuProAlaProAsnTyrThrPheAlaLeuTrpArgValSerAlaGluGluTyr  
4861 GTACCCCCCTTCTGCGCCGAACCTACAGTTCGCGCTATGGAGGGTGTCTGCAGAGGAAT  
CATGGGGGAAGGACGCGGCTTGATGTCAAGCGGATACCTCCACAGACGTCTCCTTA

FIG. 47-5

ValGluIleArgGlnValGlyAspPheHisTyrValThrGlyMetThrThrAspAsnLeu  
 4921 ATGTGGAGATAAGGCAGGTGGGGGACTTCCACTACGTGACGGGTATGACTACTGACAATC  
 TACACCTCTATTCCGTCCACCCCTGAAGGTGATGCACCTGCCATACTGATGACTGTTAG

LysCysProCysGlnValProSerProGluPhePheThrGluLeuAspGlyValArgLeu  
 4981 TCAAATGCCCGTGCCAGGTCCCATCGCCCGAATTTTTCACAGAATTGGACGGGGTGGCGC  
 AGTTTACGGGCACGGTCCAGGTAGCGGGCTTAAAAAGTGCTTAACCTGCCCCACGCGG

HisArgPheAlaProProCysLysProLeuLeuArgGluGluValSerPheArgValGly  
 5041 TACATAGGTTTGGCCCCCTGCAAGCCCTTGCTGCGGGAGGAGGTATCATTGAGAGTAG  
 ATGATATCCAAACGCGGGGGGACGTTCGGGAACGACGCCCTCTCCATAGTAAGTCTCATC

LeuHisGluTyrProValGlySerGlnLeuProCysGluProGluProAspValAlaVal  
 5101 GACTCCACGAATACCCGGTAGGGTCGCAATTACCTTGCGAGCCCGAACCGGACGTGGCCG  
 CTGAGGTGCTTATGGGCCATCCAGCGTTAATGGAACGCTCGGGCTTGGCTGCACCGCG

LeuThrSerMetLeuThrAspProSerHisIleThrAlaGluAlaAlaGlyArgArgLeu  
 5161 TGTGTAGCTCCATGCTCACTGATCCCTCCCATATAACAGCAGAGGCGGGCGGGCGAAGGT  
 ACAACTGCAGGTACGAGTACTAGGGAGGGTATATTGTCGTCTCCGCCGGCCCGCTTCCA

AlaArgGlySerProProSerValAlaSerSerSerAlaSerGlnLeuSerAlaProSer  
 5221 TGGCGAGGGGATCACCCCTCTGTGGCCAGCTCCTCGGCTAGCCAGCTATCCGCTCCAT  
 ACCGCTCCCTAGTGGGGGGAGACACCGGTGAGGAGCCGATCGGTGATAGGCGAGGTA

LeuLysAlaThrCysThrAlaAsnHisAspSerProAspAlaGluLeuIleGluAlaAsn  
 5281 CTCTCAAGCAACTTGCACCGCTAACCATGACTCCCTGATGCTGAGCTCATAGAGGCCA  
 GAGAGTTCGTTGAACGTGGCGATTGGTACTGAGGGGACTACGACTCGAGTATCTCCGGT

LeuLeuTrpArgGlnGluMetGlyGlyAsnIleThrArgValGluSerGluAsnLysVal  
 5341 ACCTCTATGGAGGCAGGAGATGGGCGGCAACATCACCAGGGTTGAGTCAGAAAACAAG  
 TGGAGGATACCTCCGTCTCTACCCGCCGTGTAGTGGTCCCAACTCAGTCTTTTGTTC

ValIleLeuAspSerPheAspProLeuValAlaGluGluAspGluArgGluIleSerVal  
 5401 TGGTGATTCTGGACTCCTTCGATCCGCTTGTTGGCGGAGGAGACGAGCGGGAGATCTCCG  
 ACCACTAAGACCTGAGGAAGCTAGGCGAACACCGCCTCTCTGCTCGCCCTCTAGAGGC

ProAlaGluIleLeuArgLysSerArgArgPheAlaGlnAlaLeuProValTrpAlaArg  
 5461 TACCCGAGAAATCTGCGGAAGTCTCGGAGATTGCGCCAGGCCCTGCGCGTTTGGCGCG  
 ATGGGCGTCTTTAGGACGCCCTCAGAGCCTTAAGCGGGTCCGGGACGGGCAAACCCGCG

ProAspTyrAsnProProLeuValGluThrTrpLysLysProAspTyrGluProProVal  
 5521 GGCCGGACTATAACCCCGCTAGTGGAGACGTGGAAAAAGCCGACTACGAACCACCTG  
 CCGCCTGATATTGGGGGGGATCACCTCTGCACCTTTTTCGGGCTGATGCTTGGTGGAC

ValHisGlyCysProLeuProProProLysSerProProValProProProArgLysLys  
 5581 TGGTCCATGGCTGTCCGCTTCCACCTCAAAGTCCCTCTGTGCTCCGCTCGGAAGA  
 ACCAGGTACCGACAGGCGAAGGTGGAGGTTTCAGGGGAGGACACGGAGGCGGAGCCTTCT

ArgThrValValLeuThrGluSerThrLeuSerThrAlaLeuAlaGluLeuAlaThrArg  
 5641 AGCGGACGGTGGTCTCTACTGAATCAACCTATCTACTGCCTTGGCCGAGCTCGCCACCA  
 TCGCTGCCACAGGAGTACTTAGTGGGATAGATGACGGAACCGGCTCGAGCGGTGGT

SerPheGlySerSerSerThrSerGlyIleThrGlyAspAsnThrThrThrSerSerGlu  
 5701 GAAGCTTTGGCAGCTCCTCAACTTCCGGCATTACGGGCGACAATACGACAACATCCTCTG  
 CTTGAAACCGTCGAGGAGTTGAAGGCCGTAATGCCCGCTGTTATGCTGTTGTAGGAGAC

ProAlaProSerGlyCysProProAspSerAspAlaGluSerTyrSerSerMetProPro  
 5761 AGCCCGCCCTTCTGGCTGCCCCCGACTCCGACGCTGAGTCTTATCTCCATGCCCC  
 TCGGGCGGGGAAGACCGACGGGGGGGCTGAGGCTGCGACTCAGGATAAGGAGGTACGGG

LeuGluGlyGluProGlyAspProAspLeuSerAspGlySerTrpSerThrValSerSer  
 5821 CCCTGGAGGGGGAGCCTGGGGATCCGGATCTTAGCGACGGGTGATGGTCAACGGTCAGTA  
 GGGACCTCCCCCTCGGACCCCTAGGCCTAGAATCGCTGCCAGTACCAGTTGCCAGTCAT

GluAlaAsnAlaGluAspValValCysCysSerMetSerTyrSerTrpThrGlyAlaLeu

FIG. 47-6

5881 GTGAGGCCAACGCGGAGGATGTCGTGTGCTGCTCAATGTCTTACTCTTGGACAGGCGCAC  
 CACTCCGGTTGCGCCTCTACAGCACACGACGAGTTACAGAATGAGAACCTGTCCGCGTG  
  
 ValThrProCysAlaAlaGluGluGlnLysLeuProIleAsnAlaLeuSerAsnSerLeu  
 5941 TCGTACCCCCGTGCGCCGCGGAAGAACAGAACTGCCATCAATGCCTAAGCAACTCGT  
 AGCAGTGGGGCACGCGCGCCTTCTTGTCTTTGACGGGTAGTTACGTGATTCTGTTGAGCA  
  
 LeuArgHisHisAsnLeuValTyrSerThrThrSerArgSerAlaCysGlnArgGlnLys  
 6001 TGCTACGTACCACAATTGGTGTATTCCACCACCTCACGAGTGTGTCCTGCAAAGGCAGA  
 ACATGCAGTGGTGTAAACCACATAAGGTGGTGGAGTGGTGCACGAACGGTTTCCGTCT  
  
 LysValThrPheAspArgLeuGlnValLeuAspSerHisTyrGlnAspValLeuLysGlu  
 6061 AGAAAGTCACATTGACAGACTGCAAGTTCGACAGCCATTACCAGGACGTACTCAAGG  
 TCTTTCAGTGTAACGTCTGACGTTCAAGACCTGTCGGTAATGGTCTGTCATGAGTTCC  
  
 ValLysAlaAlaAlaSerLysValLysAlaAsnLeuLeuSerValGluGluAlaCysSer  
 6121 AGGTTAAAGCAGCGCGCTCAAAAGTGAAGGCTAACTTGCTATCCGTAGAGGAAGCTTGCA  
 TCCAATTTCTGTCGCCGCGAGTTTTCACCTCCGATTGAACGATAGGCATCTCCTTCGAACGT  
  
 LeuThrProProHisSerAlaLysSerLysPheGlyTyrGlyAlaLysAspValArgCys  
 6181 GCCTGACGCCCCACACTCAGCCAAATCCAAGTTTGGTTATGGGGCAAAGACGTCCGTT  
 CGGACTGCGGGGGTGTGAGTCGGTTTAGGTTCAAACCAATACCCGTTTCTGTCAGGCAA  
  
 HisAlaArgLysAlaValThrHisIleAsnSerValTrpLysAspLeuLeuGluAspAsn  
 6241 GCCATGCCAGAAAGGCCGTAAACCCACATCAACTCCGTGTGGAAAGACCTTCTGGAAGACA  
 CGGTACGGTCTTTCGCGCATGGGTGTAGTTGAGGCACACCTTCTGGAAGACCTTCTGT  
  
 ValThrProIleAspThrThrIleMetAlaLysAsnGluValPheCysValGlnProGlu  
 6301 ATGTAAACCAATAGACACTACCATCATGGCTAAGAACGAGGTTTTCTGCGTTCAGCCTG  
 TACATTGTGGTTATCTGTGATGGTAGTACCGATTCTTGCTCCAAAGACGCAAGTCGGAC  
  
 LysGlyGlyArgLysProAlaArgLeuIleValPheProAspLeuGlyValArgValCys  
 6361 AGAAGGGGGGTGCTAAGCCAGCTCGTCTCATCGTGTTCCTCCGATCTGGGCGTGCGCGTGT  
 TCTTCCCCCAGCATTCCGTGAGCAGAGTAGCACAAGGGGCTAGACCCGCACGCGCACA  
  
 GluLysMetAlaLeuTyrAspValValThrLysLeuProLeuAlaValMetGlySerSer  
 6421 GCGAAAAGATGGCTTTGTACGACGTGGTTACAAAGCTCCCTTGGCCGTGATGGGAAGCT  
 CGCTTTTCTACCGAAACATGCTGCACCAATGTTTCGAGGGGAACCGGCACTACCTTCGA  
  
 TyrGlyPheGlnTyrSerProGlyGlnArgValGluPheLeuValGlnAlaTrpLysSer  
 6481 CCTACGGATTCCAATACTCACCAGACAGCGGGTTGAATTCCTCGTGCAAGCGTGGAAGT  
 GGATGCCAAGGTTATGAGTGGTCTGTCGCCCACTTAAGGAGCACGTTCCGACCTTCA  
  
 LysLysThrProMetGlyPheSerTyrAspThrArgCysPheAspSerThrValThrGlu  
 6541 CCAAGAAAACCCCAATGGGGTCTCGTATGATACCCGCTGCTTGACTCCACAGTCACTG  
 GGTCTTTTGGGGTTACCCCAAGAGCATACTATGGGCGACGAAACTGAGGTGTCAGTGAC  
  
 SerAspIleArgThrGluGluAlaIleTyrGlnCysCysAspLeuAspProGlnAlaArg  
 6601 AGAGCGACATCCGTACGGAGGAGGCAATCTACCAATGTTGTGACCTCGACCCCCAAGCCC  
 TCTCGCTGTAGGCATGCCTCCTCCGTTAGATGGTTACAACACTGGAGCTGGGGGTTCCGG  
  
 ValAlaIleLysSerLeuThrGluArgLeuTyrValGlyGlyProLeuThrAsnSerArg  
 6661 GCGTGGCATCAAGTCCCTCACCAGAGAGGCTTATGTTGGGGGCCCTCTTACCAATTCAA  
 CGCACCGGTAGTTCAGGGAGTGGCTCTCCGAAATACAACCCCGGGAGAATGGTTAAGTT  
  
 GlyGluAsnCysGlyTyrArgArgCysArgAlaSerGlyValLeuThrThrSerCysGly  
 6721 GGGGGGAGAACTGCGGCTATCGCAGGTGCGCGCGAGCGGCTACTGACAACTAGCTGTG  
 CCCCCCTTTGACGCCGATAGCGTCCACGGCGCGCTCGCCGATGACTGTTGATCGACAC  
  
 AsnThrLeuThrCysTyrIleLysAlaArgAlaAlaCysArgAlaAlaGlyLeuGlnAsp  
 6781 GTAACACCTCACTTGCTACATCAAGGCCGGGCAGCCTGTGAGCCGCGAGGGCTCCAGG  
 CATGTGGGAGTGAACGATGTAGTTCCGGGCCCGTCCGACAGCTCGGCGTCCCGAGGTCC  
  
 CysThrMetLeuValCysGlyAspAspLeuValValIleCysGluSerAlaGlyValGln  
 6841 ACTGCACCATGCTCGTGTGTGGCGACGACTAGTCGTTATCTGTGAAAGCGCGGGGGTCC  
 TGACGTGGTACGAGCACACACCGCTGCTGAATCAGCAATAGACACTTTCGCGCCCCCAGG

FIG. 47-7



6901      GluAspAlaAlaSerLeuArgAlaPheThrGluAlaMetThrArgTyrSerAlaProPro  
 AGGAGGACGCGGCGAGCCTGAGAGCCTTCACGGAGGCTATGACCAGGTACTCCGCCCCC  
 TCCTCCTGCGCGCTCGGACTCTCGGAAGTGCTCCGATACTGGTCCATGAGGCGGGGG

6961      GlyAspProProGlnProGluTyrAspLeuGluLeuIleThrSerCysSerSerAsnVal  
 CTGGGGACCCCCACAACCAGAATACGACTTGGAGCTCATAACATCATGCTCCTCCAACG  
 GACCCCTGGGGGGTGTGGTCTTATGCTGAACCTCGAGTATTGTAGTACGAGGAGGTTC

7021      SerValAlaHisAspGlyAlaGlyLysArgValTyrTyrLeuThrArgAspProThrThr  
 TGTCAGTCGCCCACGACGGCGCTGGAAAGAGGGTCTACTACCTACCCGTGACCCACAA  
 ACAGTCAGCGGGTGCTGCCGCGACCTTTCTCCAGATGATGGAGTGGGCACTGGGATGTT

7081      ProLeuAlaArgAlaAlaTrpGluThrAlaArgHisThrProValAsnSerTrpLeuGly  
 CCCCCCTCGCGAGAGCTGCGTGGGAGACAGCAAGACACACTCCAGTCAATTCCTGGCTAG  
 GGGGGGAGCGCTCTCGACGCACCCTCTGTCGTTCTGTGTGAGGTCAGTTAAGGACCGATC

7141      AsnIleIleMetPheAlaProThrLeuTrpAlaArgMetIleLeuMetThrHisPhePhe  
 GCAACATAATCATGTTTGCCCCACACTGTGGGCGAGGATGATACTGATGACCCATTCT  
 CGTGTATTAGTACAAACGGGGGTGTGACACCCGCTCCTACTATGACTACTGGGTAAAGA

7201      SerValLeuIleAlaArgAspGlnLeuGluGlnAlaLeuAspCysGluIleTyrGlyAla  
 TTAGCGTCTTATAGCCAGGGACAGCTTGAACAGGCCCTCGATTGCGAGATCTACGGGG  
 AATCGCAGGAATATCGGTCCCTGGTGAACCTGTCCGGGAGCTAACGCTCTAGATGCCCC

7261      CysTyrSerIleGluProLeuAspLeuProProIleIleGlnArgLeu  
 CCTGCTACTCCATAGAACCCTTGATCTACCTCCAATCATTCAAAGACTC  
 GGACGATGAGGTATCTTGGTGAACCTAGATGGAGGTTAGTAAGTTTCTGAG

FIG. 47- 8